SEQUENCE LISTING

<110> CHUGAI RESEARCH INSTITUTE FOR MOLECULAR MEDICINE, INC.

<120> Tumor suppressor gene

<130> C1-104PCT

<140>

<141>

<150> JP 1999-118806

<151> 1999-04-26

<160> 15

<170> PatentIn Ver. 2.0

<210> 1

<211> 2829

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (52).. (1140)

<4	Λ	۸١	. 1	
\4	ĿV	v	1	

ctccggcctt	ggtggcgggt	ggctggcggt	tccgttaggt	ctgagggagc	g	atg	gcg	57
						Met	Ala	

1

gta	cgc	gcg	ttg	aag	ctg	ctg	acc	aca	ctg	ctg	gct	gtc	gtg	gcc	gct	105
Val	Arg	Ala	Leu	Lys	Leu	Leu	Thr	Thr	Leu	Leu	Ala	Val	Val	Ala	Ala	
		5					10					15				

gcc tcc caa gcc gag gtc gag tcc gag gca gga tgg ggc atg gtg acg 153

Ala Ser Gln Ala Glu Val Glu Ser Glu Ala Gly Trp Gly Met Val Thr

20 25 30

cct gat ctg ctc ttc gcc gag ggg acc gca gcc tac gcg cgc ggg gac 201
Pro Asp Leu Leu Phe Ala Glu Gly Thr Ala Ala Tyr Ala Arg Gly Asp

40
45
50

tgg ccc ggg gtg gtc ctg agc atg gaa cgg gcg ctg cgc tcc cgg gca 249

Trp Pro Gly Val Val Leu Ser Met Glu Arg Ala Leu Arg Ser Arg Ala

55 60 65

gcc ctc cgc gcc ctt cgc ctg cgc tgc cgc acc cag tgt gcc gcc gac 297

Ala Leu Arg Ala Leu Arg Leu Arg Cys Arg Thr Gln Cys Ala Ala Asp

70 75 80

ttc	ccg	tgg	gag	ctg	gac	ccc	gac	tgg	tcc	ccc	agc	ccg	gcc	cag	gcc	345
Phe	Pro	Trp	Glu	Leu	Asp	Pro	Asp	Trp	Ser	Pro	Ser	Pro	Ala	Gln	Ala	
		85					90					95				
tcg	ggc	gcc	ggc	gcc	ctg	cgc	gac	ctg	agc	ttc	ttc	ggg	ggc	ctt	ctg	393
Ser	Gly	Ala	Gly	Ala	Leu	Arg	Asp	Leu	Ser	Phe	Phe	Gly	Gly	Leu	Leu	
	100					105					110					
cgt	cgc	gct	gcc	tgc	ctg	cgc	cgc	tgc	ctc	ggg	ccg	ccg	gcc	gcc	cac	441
Arg	Arg	Ala	Ala	Cys	Leu	Arg	Arg	Cys	Leu	Gly	Pro	Pro	Ala	Ala	His	
115					120					125					130	
tcg	ctc	agc	gaa	gag	atg	gag	ctg	gag	ttc	cgc	aag	cgg	agc	ccc	tac	489
Ser	Leu	Ser	Glu	Glu	Met	Glu	Leu	Glu	Phe	Arg	Lys	Arg	Ser	Pro	Tyr	
				135					140					145		
											_					
aac	tac	ctg	cag	gtc	gcc	tac	ttc	aag	atc	aac	aag	ttg	gag	aaa	gct	537
Asn	Tyr	Leu	Gln	Val	Ala	Tyr	Phe	Lys	Ile	Asn	Lys	Leu	Glu	Lys	Ala	
			150					155					160			
gtt	gct	gca	gca	cac	acc	ttc	ttc	gtg	ggc	aat	cct	gag	cac	atg	gaa	585
/al	Ala	Ala	Ala	His	Thr	Phe	Phe	Val	Gly	Asn	Pro	Glu	His	Met	Glu	
		165					170					175				

atg cag cag aac cta gac tat tac caa acc atg tct gga gtg aag gag 633 Met Gln Gln Asn Leu Asp Tyr Tyr Gln Thr Met Ser Gly Val Lys Glu

The street street is the street stree

gco	gac	ttc	aag	g gat	ctt	gag	act	caa	ccc	cat	ate	; caa	gaa	ttt	cga	681
Ala	a Asp	Phe	Lys	s Asp	Leu	Glu	Thr	Gln	Pro	His	Met	Gln	Glu	. Phe	Arg	
195	5				200					205	5				210	
ctg	gga	gtg	cga	ctc	tac	tca	gag	gaa	cag	cca	cag	gaa	gct	gtg	ccc	729
Leu	Gly	Val	Arg	Leu	Tyr	Ser	Glu	Glu	Gln	Pro	Gln	Glu	Ala	. Val	Pro	
				215					220					225		
cac	cta	gag	gcg	gcg	ctg	caa	gaa	tac	ttt	gtg	gcc	tat	gag	gag	tgc	777
	Leu															
			230					235				-	240		-	
cgt	gcc	ctc	tgc	gaa	ggg	ccc	tat	gac	tac	gat	ggc	tac	aac	tac	ctt	825
	Ala															
		245					250			•	-	255		- ,		
gag	tac	aac	gct	gac	ctc	ttc	cag	gcc	atc	aca	gat.	cat	tac	atc	cag	873
	Tyr															0.0
	260			-		265					270	1110	1,1	110	OIII	
											210					
gtc	ctc	aac	tgt	aag	cag	aac	t.gt.	gtc	മറത	asa	ctt	act	too	000	000	Q91
	Leu															921
			- , -	_, ~			J J U	. 41	TIIT	JIU	Leu	nia	DGI.	IITZ	LIO	

agt cga gag aag ccc ttt gaa gac ttc ctc cca tcg cat tat aat tat 969

Ser Arg Glu Lys Pro Phe Glu Asp Phe Leu Pro Ser His Tyr Asn Tyr

295 300 305

ctg cag ttt gcc tac tat aac att ggg aat tat aca caa gct ggt gaa 1017 Leu Gln Phe Ala Tyr Tyr Asn Ile Gly Asn Tyr Thr Gln Ala Gly Glu 310 315 320

tgt gcc aag acc tat ctt ctc ttc ttc ccc aat gac gag gtg atg aac 1065

Cys Ala Lys Thr Tyr Leu Leu Phe Phe Pro Asn Asp Glu Val Met Asn

325 330 335

caa aat ttg gcc tat tat gca gct atg ctt gga gaa gaa cac acc aga 1113 Gln Asn Leu Ala Tyr Tyr Ala Ala Met Leu Gly Glu Glu His Thr Arg 340 345 350

tcc atc ggc ccc cgt gag cag ggc acc tagggaaaga tgtgaccccg

1160

Ser Ile Gly Pro Arg Glu Gln Gly Thr

355

360

gaaagtactc agtttccctg ccctggagtg ccaaggagta ccgacagcga agcctactgg 1220

aaaaagaact gcttttcttc gcttatgatg tttttggaat tccctttgtg gatcgggatt 1280

catggactcc agaagaaatg attcccaaga aattgcaaga gaaacagaag tgaggacctt 1340

gaagaaactg catggttgga tcagtctgat gaagcacttg aggcttcttg agcccaggca 1400 gatgtgaact cctggcaagg ggtgggcagg tccagtttgg gaagtcgggg tggagcccag 1460 ggctggccct ggaatgcagt cctcagagcg gttgtgctca taggtcagaa cgggaaacag 1520 ccgtacgcat ctcccaggag attgggaacc ttatgaagga aatcgagacc cttgtggaag 1580 agaagaccaa ggagtcactg gatgtgagca gactgacccg ggaaggtggc cccctgctgt 1640 atgaaggcat cagtctcacc atgaactcca aactcctgaa tggttaccag cgggtggtga 1700 tggacggcgt aatctctgac cacgagtgtc aggagctgca gagactgacc aatgtggcag 1760 caacctcagg agatggctac cggggtcaga cctccccaca tactcccaat gaaaagttct 1820 atggtgtcac tgtcttcaaa gccctcaagc tggggcaaga aggcaaagtt cctctgcaga 1880 gtgcccacct gtactacaac gtgacggaga aagtgcggcg catcatggag tcctacttcc 1940 gcctggatac gccctctac ttttcctact ctcatctggt gtgccgcact gccatcgaag 2000 aggtccaggc agagaggaag gatgatagtc atccagtcca cgtggacaac tgcatcctga 2060 atgccgagac cctcgtgtgt gtcaaagagc ccccagccta caccttccgc gactacagcg 2120

ccatccttta cctaaatggg gacttcgatg gcggaaactt ttatttcact gaactggatg 2180 ccaagaccgt gacggcagag gtgcagcctc agtgtggaag agccgtggga ttctcttcag 2240 gcactgaaaa cccacatgga gtgaaggctg tcaccagggg gcagcgctgt gccatcgccc 2300 tgtggttcac cctggaccct cgacacagcg agcgggacag ggtgcaggca gatgacctgg 2360 tgaagatgct cttcagccca gaagagatgg acctctccca ggagcagccc ctggatgccc 2420 agcagggccc ccccgaacct gcacaagagt ctctctcagg cagtgaatcg aagcccaagg 2480 atgagctatg acagcgtcca ggtcagacgg atgggtgact agacccatga agaggaactc 2540 ttcttgcact ctgagctggc cagcccctcg gggctgcaga gcagtgagcc tacatctgcc 2600 actcagccga ggggaccctg ctcacagcct tctacatggt gctactgctc ttggagtgga 2660 catgaccaga caccgcaccc cctggatctg gctgagggct caggacacag gcccagccac 2720 ccccagggc ctccacaggc cgctgcataa cagcgataca gtacttaagt gtctgtgtag 2780 acaaccaaag aataaatgat tcatggtttt ttttaaaaaa aaaaaaaaa 2829

<211> 363 <212> PRT

<213> Homo sapiens

<400> 2

Met Ala Val Arg Ala Leu Lys Leu Leu Thr Thr Leu Leu Ala Val Val

1 5 10 15

Ala Ala Ser Gln Ala Glu Val Glu Ser Glu Ala Gly Trp Gly Met
20 25 30

Val Thr Pro Asp Leu Leu Phe Ala Glu Gly Thr Ala Ala Tyr Ala Arg

35 40 45

Gly Asp Trp Pro Gly Val Val Leu Ser Met Glu Arg Ala Leu Arg Ser 50 55 60

Arg Ala Ala Leu Arg Ala Leu Arg Leu Arg Cys Arg Thr Gln Cys Ala 65 70 75 80

Ala Asp Phe Pro Trp Glu Leu Asp Pro Asp Trp Ser Pro Ser Pro Ala
85 90 95

Gln Ala Ser Gly Ala Gly Ala Leu Arg Asp Leu Ser Phe Phe Gly Gly
100 105 110

Leu Leu Arg Arg Ala Ala Cys Leu Arg Arg Cys Leu Gly Pro Pro Ala
115 120 125

Ala His Ser Leu Ser Glu Glu Met Glu Leu Glu Phe Arg Lys Arg Ser

130 135 140

Pro Tyr Asn Tyr Leu Gln Val Ala Tyr Phe Lys Ile Asn Lys Leu Glu
145 150 155 160

Lys Ala Val Ala Ala Ala His Thr Phe Phe Val Gly Asn Pro Glu His
165 170 175

Met Glu Met Gln Gln Asn Leu Asp Tyr Tyr Gln Thr Met Ser Gly Val
180 185 190

Lys Glu Ala Asp Phe Lys Asp Leu Glu Thr Gln Pro His Met Gln Glu
195 200 205

Phe Arg Leu Gly Val Arg Leu Tyr Ser Glu Glu Gln Pro Gln Glu Ala 210 215 220

Val Pro His Leu Glu Ala Ala Leu Gln Glu Tyr Phe Val Ala Tyr Glu 225 230 235 240

Glu Cys Arg Ala Leu Cys Glu Gly Pro Tyr Asp Tyr Asp Gly Tyr Asn 245 250 255 Tyr Leu Glu Tyr Asn Ala Asp Leu Phe Gln Ala Ile Thr Asp His Tyr
260 265 270

Ile Gln Val Leu Asn Cys Lys Gln Asn Cys Val Thr Glu Leu Ala Ser 275 280 285

His Pro Ser Arg Glu Lys Pro Phe Glu Asp Phe Leu Pro Ser His Tyr
290 295 300

Asn Tyr Leu Gln Phe Ala Tyr Tyr Asn Ile Gly Asn Tyr Thr Gln Ala 305 310 315 320

Gly Glu Cys Ala Lys Thr Tyr Leu Leu Phe Phe Pro Asn Asp Glu Val
325 330 335

Met Asn Gln Asn Leu Ala Tyr Tyr Ala Ala Met Leu Gly Glu Glu His

340 345 350

Thr Arg Ser Ile Gly Pro Arg Glu Gln Gly Thr
355 360

⟨210⟩ 3

<211> 2600

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (52).. (2259)

<400> 3

ctccggcctt ggtggcggt ggctggcggt tccgttaggt ctgagggagc g atg gcg 57

Met Ala

1

gta cgc gcg ttg aag ctg ctg acc aca ctg ctg gct gtc gtg gcc gct 105

Val Arg Ala Leu Lys Leu Leu Thr Thr Leu Leu Ala Val Val Ala Ala

5 10 15

gcc tcc caa gcc gag gtc gag tcc gag gca gga tgg ggc atg gtg acg 153

Ala Ser Gln Ala Glu Val Glu Ser Glu Ala Gly Trp Gly Met Val Thr

20 25 30

cct gat ctg ctc ttc gcc gag ggg acc gca gcc tac gcg cgc ggg gac 201
Pro Asp Leu Leu Phe Ala Glu Gly Thr Ala Ala Tyr Ala Arg Gly Asp
35 40 45 50

tgg ccc ggg gtg gtc ctg agc atg gaa cgg gcg ctg cgc tcc cgg gca 249

Trp Pro Gly Val Val Leu Ser Met Glu Arg Ala Leu Arg Ser Arg Ala

55 60 65

297	gac	gcc	gcc	tgt	cag	acc	cgc	tgc	cgc	ctg	cgc	ctt	gcc	cgc	ctc	gcc
	Asp	Ala	Ala	Cys	Gln	Thr	Arg	Cys	Arg	Leu	Arg	Leu	Ala	Arg	Leu	Ala
			80					75					70			
345	gcc	cag	gcc	ccg	agc	ccc	tcc	tgg	gac	ccc	gac	ctg	gag	tgg	ccg	ttc
	Ala	Gln	Ala	Pro	Ser	Pro	Ser	Trp	Asp	Pro	Asp	Leu	Glu	Trp	Pro	Phe
				95					90					85		
393	ctg	ctt	ggc	ggg	ttc	ttc	agc	ctg	gac	cgc	ctg	gcc	ggc	gcc	ggc	tcg
	Leu	Leu	Gly	G1y	Phe	Phe	Ser	Leu	Asp	Arg	Leu	Ala	Gly	Ala	Gly	Ser
					110					105					100	
441	cac	gcc	gcc	ccg	ccg	ggg	ctc	tgc	cgc	cgc	ctg	tgc	gcc	gct	cgc	cgt
	His	Ala	Ala	Pro	Pro	G1y	Leu	Cys	Arg	Arg	Leu	Cys	Ala	Ala	Arg	Arg
	130					125					120					115
					-											
489	tac	ccc	agc	cgg	aag	cgc	ttc	gag	ctg	gag	atg	gag	gaa	agc	ctc	tcg
	Tyr	Pro	Ser	Arg	Lys	Arg	Phe	Glu	Leu	Glu	Met	Glu	Glu	Ser	Leu	Ser
		145					140					135				
537	gct	aaa	gag	ttg	aag	aac	atc	aag	ttc	tac	gcc	gtc	cag	ctg	tac	aac

gtt gct gca gca cac acc ttc ttc gtg ggc aat cct gag cac atg gaa 585

155

160

Asn Tyr Leu Gln Val Ala Tyr Phe Lys Ile Asn Lys Leu Glu Lys Ala

150

Val	Ala	Ala	Ala	His	Thr	Phe	Phe	Val	Gly	Asn	Pro	Glu	His	Met	Glu	
		165					170					175				
ato	cag	cag	aac	cta	gac	tat	tac	caa	acc	ato	tet	aaa	at a	221	as a	633
				_		_										000
Met	Gln	Gln	Asn	Leu	Asp	Tyr	Tyr	Gln	Thr	Met	Ser	Gly	Val	Lys	Glu	
	180					185					190					
gcc	gac	ttc	aag	gat	ctt	gag	act	caa	ccc	cat	atg	caa	gaa	ttt	cga	681
Ala	Asp	Phe	Lys	Asp	Leu	Glu	Thr	G1n	Pro	His	Met	Gln	Glu	Phe	Arg	
195					200					205					210	
ctg	gga	gtg	cga	ctc	tac	tca	gag	gaa	cag	cca	cag	ฮลล	gct	o to	ccc	729
																120
Leu	Gly	vaı	Arg		ıyr	ser	GIU	GIU		Pro	Gin	Glu	Ala	Vai	Pro	
				215					220					225		
cac	cta	gag	gcg	gcg	ctg	caa	gaa	tac	ttt	gtg	gcc	tat	gag	gag	tgc	777
His	Leu	Glu	Ala	Ala	Leu	G1n	Glu	Tyr	Phe	Val	Ala	Tyr	Glu	G1u	Cys	
			230					235					240			
cøt	gcc	ctc	t.gc	gaa	ggg	ccc	tat	gan	tac	as t	aac	tan	220	taa	at+	825
																040
Arg	Ala	Leu	Cys	Glu	Gly	Pro	Tyr	Asp	Tyr	Asp	Gly	Tyr	Asn	Tyr	Leu	

gag tac aac gct gac ctc ttc cag gcc atc aca gat cat tac atc cag 873 Glu Tyr Asn Ala Asp Leu Phe Gln Ala Ile Thr Asp His Tyr Ile Gln

255

250

260 265 270

245

gtc	ctc	aac	tgt	aag	cag	aac	tgt	gtc	acg	gag	cţt	gct	tcc	cac	cca	921
Val	Leu	Asn	Cys	Lys	G1n	Asn	Cys	Val	Thr	Glu	Leu	Ala	Ser	His	Pro	
275					280					285					290	
agt	cga	gag	aag	ccc	ttt	gaa	gac	ttc	ctc	cca	tcg	cat	tat	aat	tat	969
	Arg															
261	Λιg	Giu	Lys		1 116	Giu	лър	1 116		110	per	1112	1 1 1		1 9 1	
				295					300					305		
ctg	cag	ttt	gcc	tac	tat	aac	att	ggg	aat	tat	aca	caa	gct	ggt	gaa	1017
Leu	Gln	Phe	Ala	Tyr	Tyr	Asn	Ile	G1y	Asn	Tyr	Thr	G1n	Ala	Gly	Glu	
			310					315					320			
tgt	gcc	aag	acc	tat	ctt	ctc	ttc	ttc	ccc	aat.	gac	១ឧទ	gt.g	atø	aac	1065
	Ala															1000
Cys	nia		1111	lyı	Leu	Leu		rne	110	ASII	ASP		Val	Met	ASII	
		325					330				-	335				
caa	aat	ttg	gcc	tat	tat	gca	gct	atg	ctt	gga	gaa	gaa	cac	acc	aga	1113
Gln	Asn	Leu	Ala	Tyr	Tyr	Ala	Ala	Met	Leu	Gly	Glu	Glu	His	Thr	Arg	
	340					345					350					
tcc	atc	ggc	ccc	cgt	gag	agt.	gcc	ลลด	ទួនថ	tac	റമ്മ	cag	COS	age	cta	1161
																1101
	Ile	ату	LTO	VT.R		net.	ита	LyS	GIU		Arg	GIN	arg	ser		
355					360					365					370	

ctg gaa aaa gaa ctg ctt ttc ttc gct tat gat gtt ttt gga att ccc 1209

Leu	Glu	Lys	Glu	Leu	Leu	Phe	Phe	Ala	Tyr	Asp	Val	Phe	Gly	Ile	Pro
				375					380		-			385	

ttt gtg gat ccg gat tca tgg act cca gaa gaa gtg att ccc aag aga 1257
Phe Val Asp Pro Asp Ser Trp Thr Pro Glu Glu Val Ile Pro Lys Arg
390 395 400

ttg caa gag aaa cag aag tca gaa cgg gaa aca gcc gta cgc atc tcc 1305 Leu Gln Glu Lys Gln Lys Ser Glu Arg Glu Thr Ala Val Arg Ile Ser 405 410 415

cag gag att ggg aac ctt atg aag gaa atc gag acc ctt gtg gaa gag 1353 Gln Glu Ile Gly Asn Leu Met Lys Glu Ile Glu Thr Leu Val Glu Glu 420 425 430

aag acc aag gag tca ctg gat gtg agc aga ctg acc cgg gaa ggt ggc 1401 Lys Thr Lys Glu Ser Leu Asp Val Ser Arg Leu Thr Arg Glu Gly Gly 435 440 445 450

ccc ctg ctg tat gaa ggc atc agt ctc acc atg aac tcc aaa ctc ctg1449Pro Leu Leu Tyr Glu Gly Ile Ser Leu Thr Met Asn Ser Lys Leu Leu455460465

aat ggt tac cag cgg gtg gtg atg gac ggc gta atc tct gac cac gag 1497 Asn Gly Tyr Gln Arg Val Val Met Asp Gly Val Ile Ser Asp His Glu 470 475 480 tgt cag gag ctg cag aga ctg acc aat gtg gca gca acc tca gga gat 1545 Cys Gln Glu Leu Gln Arg Leu Thr Asn Val Ala Ala Thr Ser Gly Asp 485 490 495

ggc tac cgg ggt cag acc tcc cca cat act ccc aat gaa aag ttc tat 1593

Gly Tyr Arg Gly Gln Thr Ser Pro His Thr Pro Asn Glu Lys Phe Tyr

500 505 510

ggt gtc act gtc ttc aaa gcc ctc aag ctg ggg caa gaa ggc aaa gtt 1641 Gly Val Thr Val Phe Lys Ala Leu Lys Leu Gly Gln Glu Gly Lys Val 515 520 530

cct ctg cag agt gcc cac ctg tac tac aac gtg acg gag aaa gtg cgg 1689

Pro Leu Gln Ser Ala His Leu Tyr Tyr Asn Val Thr Glu Lys Val Arg
535 540 545

cgc atc atg gag tcc tac ttc cgc ctg gat acg ccc ctc tac ttt tcc 1737

Arg Ile Met Glu Ser Tyr Phe Arg Leu Asp Thr Pro Leu Tyr Phe Ser

550 555 560

tac tct cat ctg gtg tgc cgc act gcc atc gaa gag gtc cag gca gag 1785

Tyr Ser His Leu Val Cys Arg Thr Ala Ile Glu Glu Val Gln Ala Glu

565 570 575

agg aag gat gat agt cat cca gtc cac gtg gac aac tgc atc ctg aat 1833

Arg Lys Asp Asp Ser His Pro Val His Val Asp Asn Cys Ile Leu Asn 580 585 590

gcc gag acc ctc gtg tgt gtc aaa gag ccc cca gcc tac acc ttc cgc 1881

Ala Glu Thr Leu Val Cys Val Lys Glu Pro Pro Ala Tyr Thr Phe Arg

595 600 605 610

gac tac agc gcc atc ctt tac cta aat ggg gac ttc gat ggc gga aac 1929
Asp Tyr Ser Ala Ile Leu Tyr Leu Asn Gly Asp Phe Asp Gly Gly Asn
615 620 625

ttt tat ttc act gaa ctg gat gcc aag acc gtg acg gca gag gtg cag 1977

Phe Tyr Phe Thr Glu Leu Asp Ala Lys Thr Val Thr Ala Glu Val Gln
630 635 640

cct cag tgt gga aga gcc gtg gga ttc tct tca ggc act gaa aac cca 2025 Pro Gln Cys Gly Arg Ala Val Gly Phe Ser Ser Gly Thr Glu Asn Pro 645 650 655

cat gga gtg aag gct gtc acc agg ggg cag cgc tgt gcc atc gcc ctg 2073

His Gly Val Lys Ala Val Thr Arg Gly Gln Arg Cys Ala Ile Ala Leu

660 665 670

tgg ttc acc ctg gac cct cga cac agc gag cgg gac agg gtg cag gca 2121

Trp Phe Thr Leu Asp Pro Arg His Ser Glu Arg Asp Arg Val Gln Ala

675 680 685 690

				695					700				_	705		
Asp	Asp	Leu	Val	Lys	Met	Leu	Phe	Ser	Pro	Glu	Glu	Met	Asp	Leu	Ser	
gat	gac	ctg	gtg	aag	atg	ctc	ttc	agc	cca	gaa	gag	atg	gac	ctc	tcc	2169

cag gag cag ccc ctg gat gcc cag cag ggc ccc ccc gaa cct gca caa 2217

Gln Glu Gln Pro Leu Asp Ala Gln Gln Gly Pro Pro Glu Pro Ala Gln

710 715 720

gag tet etc tea gge agt gaa teg aag eec aag gat gag eta 2259

Glu Ser Leu Ser Gly Ser Glu Ser Lys Pro Lys Asp Glu Leu

725 730 735

<210> 4 <211> 736 <212> PRT <213> Homo sapiens <400> 4 Met Ala Val Arg Ala Leu Lys Leu Leu Thr Thr Leu Leu Ala Val Val Ala Ala Ser Gln Ala Glu Val Glu Ser Glu Ala Gly Trp Gly Met Val Thr Pro Asp Leu Leu Phe Ala Glu Gly Thr Ala Ala Tyr Ala Arg Gly Asp Trp Pro Gly Val Val Leu Ser Met Glu Arg Ala Leu Arg Ser Arg Ala Ala Leu Arg Ala Leu Arg Leu Arg Cys Arg Thr Gln Cys Ala Ala Asp Phe Pro Trp Glu Leu Asp Pro Asp Trp Ser Pro Ser Pro Ala Gln Ala Ser Gly Ala Gly Ala Leu Arg Asp Leu Ser Phe Phe Gly Gly

Leu Leu Arg Arg Ala Ala Cys Leu Arg Arg Cys Leu Gly Pro Pro Ala 115 120 125

Ala His Ser Leu Ser Glu Glu Met Glu Leu Glu Phe Arg Lys Arg Ser

130 135 140

Pro Tyr Asn Tyr Leu Gln Val Ala Tyr Phe Lys Ile Asn Lys Leu Glu 145 150 155 160

Lys Ala Val Ala Ala Ala His Thr Phe Phe Val Gly Asn Pro Glu His

165 170 175

Met Glu Met Gln Gln Asn Leu Asp Tyr Tyr Gln Thr Met Ser Gly Val
180 185 190

Lys Glu Ala Asp Phe Lys Asp Leu Glu Thr Gln Pro His Met Gln Glu
195 200 205

Phe Arg Leu Gly Val Arg Leu Tyr Ser Glu Glu Gln Pro Gln Glu Ala 210 215 220

Val Pro His Leu Glu Ala Ala Leu Gln Glu Tyr Phe Val Ala Tyr Glu 225 230 235 240

Glu Cys Arg Ala Leu Cys Glu Gly Pro Tyr Asp Tyr Asp Gly Tyr Asn

Tyr	Leu	Glu	Tyr	Asn	Ala	Asp	Leu	Phe	Gln	Ala	Ile	Thr	Asp	His	Tyr
			260					265					270		
Ile	Gln	Val	Leu	Asn	Cys	Lys	G1n	Asn	Cys	Val	Thr	Glu	Leu	Ala	Ser
		275					280					285			

His Pro Ser Arg Glu Lys Pro Phe Glu Asp Phe Leu Pro Ser His Tyr

Asn Tyr Leu Gln Phe Ala Tyr Tyr Asn Ile Gly Asn Tyr Thr Gln Ala

Gly Glu Cys Ala Lys Thr Tyr Leu Leu Phe Phe Pro Asn Asp Glu Val

Met Asn Gln Asn Leu Ala Tyr Tyr Ala Ala Met Leu Gly Glu Glu His

Thr Arg Ser Ile Gly Pro Arg Glu Ser Ala Lys Glu Tyr Arg Gln Arg

Ser Leu Leu Glu Lys Glu Leu Leu Phe Phe Ala Tyr Asp Val Phe Gly

Ile Pro Phe Val Asp Pro Asp Ser Trp Thr Pro Glu Glu Val Ile Pro 385 390 395 400

Lys Arg Leu Gln Glu Lys Gln Lys Ser Glu Arg Glu Thr Ala Val Arg
405 410 415

Ile Ser Gln Glu Ile Gly Asn Leu Met Lys Glu Ile Glu Thr Leu Val
420 425 430

Glu Glu Lys Thr Lys Glu Ser Leu Asp Val Ser Arg Leu Thr Arg Glu
435
440
445

Gly Gly Pro Leu Leu Tyr Glu Gly Ile Ser Leu Thr Met Asn Ser Lys
450 455 460

Leu Leu Asn Gly Tyr Gln Arg Val Val Met Asp Gly Val Ile Ser Asp
465
470
475
480

His Glu Cys Gln Glu Leu Gln Arg Leu Thr Asn Val Ala Ala Thr Ser 485 490 495

Gly Asp Gly Tyr Arg Gly Gln Thr Ser Pro His Thr Pro Asn Glu Lys
500 505 510

Phe Tyr Gly Val Thr Val Phe Lys Ala Leu Lys Leu Gly Gln Glu Gly
515 520 525

Lys Val Pro Leu Gln Ser Ala His Leu Tyr Tyr Asn Val Thr Glu Lys
530 535 540

Val Arg Arg Ile Met Glu Ser Tyr Phe Arg Leu Asp Thr Pro Leu Tyr 545 550 555 560

Phe Ser Tyr Ser His Leu Val Cys Arg Thr Ala Ile Glu Glu Val Gln
565 570 575

Ala Glu Arg Lys Asp Asp Ser His Pro Val His Val Asp Asn Cys Ile
580
585
590

Leu Asn Ala Glu Thr Leu Val Cys Val Lys Glu Pro Pro Ala Tyr Thr
595 600 605

Phe Arg Asp Tyr Ser Ala Ile Leu Tyr Leu Asn Gly Asp Phe Asp Gly 610 615 620

Gly Asn Phe Tyr Phe Thr Glu Leu Asp Ala Lys Thr Val Thr Ala Glu
625 630 635 640

Val Gln Pro Gln Cys Gly Arg Ala Val Gly Phe Ser Ser Gly Thr Glu
645 650 655

Asn Pro His Gly Val Lys Ala Val Thr Arg Gly Gln Arg Cys Ala Ile

Ala Leu Trp Phe Thr Leu Asp Pro Arg His Ser Glu Arg Asp Arg Val
675 680 685

Gln Ala Asp Asp Leu Val Lys Met Leu Phe Ser Pro Glu Glu Met Asp 690 695 700

Leu Ser Gln Glu Gln Pro Leu Asp Ala Gln Gln Gly Pro Pro Glu Pro 705 710 715 720

Ala Gln Glu Ser Leu Ser Gly Ser Glu Ser Lys Pro Lys Asp Glu Leu
725 730 735

⟨210⟩ 5

<211> 2416

<212> DNA

<213> Mus musculus

<220>

<221> CDS

<222> (12).. (2252)

<400> 5

ggagcaaggc c atg gcg gtg acg aaa gga ggc tgc tgg cac gat gct agc 50

Met Ala Val Thr Lys Gly Gly Cys Trp His Asp Ala Ser 1 5 - 10

ggt	cgc	cgc	cgc	cgc	cgc	ctt	acg	ggt	tgc	ggc	gag	tct	gag	ccg	gga	98
G1y	Arg	Arg	Arg	Arg	Arg	Leu	Thr	Gly	Cys	Gly	Glu	Ser	Glu	Pro	G1y	
	15					20					25					

tgg gac gtg gca gcc cct gac ctg ctt tac gca gag ggg acc gcg gcc 146

Trp Asp Val Ala Ala Pro Asp Leu Leu Tyr Ala Glu Gly Thr Ala Ala

30 35 40 45

tac tcg cgc agg gac tgg ccc ggg gtg gtc ctg aac atg gag cgg gct 194

Tyr Ser Arg Arg Asp Trp Pro Gly Val Val Leu Asn Met Glu Arg Ala

50 55 60

ctg cgc tcg cgg gcc ctg cgt gcc ctc cgc ctg cgc tgc cgc aca 242

Leu Arg Ser Arg Ala Ala Leu Arg Ala Leu Arg Leu Arg Cys Arg Thr

65 70 75

cgc tgt gcc acc gaa ctg ccg tgg gca ccg gac ctg gat ctc ggt ccg 290

Arg Cys Ala Thr Glu Leu Pro Trp Ala Pro Asp Leu Asp Leu Gly Pro

80 85 90

gac ccc agc ctg agc cag gac ccg ggc gcc gcc gcc ctg cac gac ctg 338

Asp Pro Ser Leu Ser Gln Asp Pro Gly Ala Ala Ala Leu His Asp Leu

95 100 105

cgc	ttc	ttc	gga	gcc	gtg	ctg	cgc	cgt	gcc	gcc	tgc-	cta	cgc	cgc	tgc	386
Arg	Phe	Phe	Gly	Ala	Val	Leu	Arg	Arg	Ala	Ala	Cys	Leu	Arg	Arg	Cys	
110					115					120					125	
ctc	ggg	ccg	ccc	tct	gcc	cac	ttg	ctg	agt	gaa	gaa	ctg	gac	ctg	gag	434
Leu	G1y	Pro	Pro	Ser	Ala	His	Leu	Leu	Ser	Glu	Glu	Leu	Asp	Leu	Glu	
				130					135					140		
ttc	aac	aag	cgg	agc	ccg	tac	aac	tac	ctg	cag	gtc	gcc	tat	ttc	aag	482
Phe	Asn	Lys	Arg	Ser	Pro	Tyr	Asn	Tyr	Leu	Gln	Val	Ala	Tyr	Phe	Lys	
			145					150					155			
ata	aac	aag	ctg	gag	aaa	gct	gtg	gct	gcg	gca	cac	acc	ttc	ttt	gtg	530
Ile	Asn	Lys	Leu	G1u	Lys	Ala	Val	Ala	Ala	Ala	His	Thr	Phe	Phe	Val	
		160					165					170				
ggc	aat	cct	gag	cac	atg	gag	atg	cgg	cag	aac	ctc	gac	tat	tac	caa	578
Gly	Asn	Pro	Glu	His	Met	Glu	Met	Arg	Gln	Asn	Leu	Asp	Tyr	Tyr	Gln	
	175					180					185					
acc	atg	tct	ggg	gtg	aag	gag	gca	gac	ttc	agg	gat	ctc	gag	gcc	aag	626
Thr	Met	Ser	Gly	Val	Lys	Glu	Ala	Asp	Phe	Arg	Asp	Leu	Glu	Ala	Lys	
190					195					200					205	

ccc cat atg cat gag ttt cgg ctg ggg gta cga ctc tac tca gag gag 674

Pro	His	Met	His	Glu	Phe	Arg	Leu	Gly	Val	Arg	Leu	Tyr	Ser	Glu	Glu	
				210					215		-			220		
aag	cca	cag	gaa	gct	gtg	ccc	cac	ctg	gag	gcg	gca	ctg	caa	gag	tac	722
Lys	Pro	Gln	Glu	Ala	Val	Pro	His	Leu	Glu	Ala	Ala	Leu	Gln	Glu	Tyr	
			225					230					235			
														,		
ttt	gtg	gcc	gat	gag	gag	tgc	cgt	gcc	ctc	tgc	gaa	ggg	ccc	tat	gac	770
Phe	Val	Ala	Asp	Glu	Glu	Cys	Arg	Ala	Leu	Cys	G1u	G1y	Pro	Tyr	Asp	
		240					245					250				
tac	gac	ggc	tac	aac	tac	cta	gac	tac	agc	gct	gac	ctc	ttc	cag	gcc	818
Tyr	Asp	Gly	Tyr	Asn	Tyr	Leu	Asp	Tyr	Ser	Ala	Asp	Leu	Phe	G1n	Ala	
	255					260					265					
atc	aca	gat	cat	tac	gtc	cag	gtc	ctc	aac	tgt	aag	cag	aac	tgt	gtc	866
Ile	Thr	Asp	His	Tyr	Val	Gln	Val	Leu	Asn	Cys	Lys	Gln	Asn	Cys	Val	
270					275					280					285	
acg	gag	ctg	gct	tcc	cac	cca	agt	agg	gaa	aag	ccc	ttt	gaa	gac	ttc	914
Thr	Glu	Leu	Ala	Ser	His	Pro	Ser	Arg	Glu	Lys	Pro	Phe	Glu	Asp	Phe	
				290					295					300		
ctc	cct	tca	cac	tat	aat	tac	cta	cag	ttt	gcc	tac	tac	aac	att	ggg	962

Leu Pro Ser His Tyr Asn Tyr Leu Gln Phe Ala Tyr Tyr Asn Ile Gly

aac tat aca caa gct att gaa tgt gcc aag acc tac ctc ctc ttc ttt 1010
Asn Tyr Thr Gln Ala Ile Glu Cys Ala Lys Thr Tyr Leu Leu Phe Phe
320 325 330

ccc aat gat gag gtg atg cac cag aat ctg gct tat tac aca gcc atg 1058

Pro Asn Asp Glu Val Met His Gln Asn Leu Ala Tyr Tyr Thr Ala Met

335 340 345

ctt gga gaa gaa gag gcc agc tcc atc agc ccc agg gag aat gcc gag 1106 Leu Gly Glu Glu Glu Ala Ser Ser Ile Ser Pro Arg Glu Asn Ala Glu 350 355 360 365

gaa tac cga cgt cca aac ctg ttg gag aaa gaa ctg ctt ttc ttc gct 1154

Glu Tyr Arg Arg Pro Asn Leu Leu Glu Lys Glu Leu Leu Phe Phe Ala

370 375 380

tat gac att ttt gga att ccc ttt gtg gat ccc gat tca tgg act cca 1202

Tyr Asp Ile Phe Gly Ile Pro Phe Val Asp Pro Asp Ser Trp Thr Pro

385 390 395

gaa gaa gtg att ccc aag aga ttg caa gag aaa cag aag tct gaa cgg 1250
Glu Glu Val Ile Pro Lys Arg Leu Gln Glu Lys Gln Lys Ser Glu Arg
400 405 410

gaa aca gcc gta cgc atc tcc cag gag att ggg aac ctt atg aag gaa 1298

Glu Thr Ala Val Arg Ile Ser Gln Glu Ile Gly Asn Leu Met Lys Glu
415 420 425

atc gag acc ctt gtg gaa gag aag acc aag gag tct ctg gat gtg agc 1346

Ile Glu Thr Leu Val Glu Glu Lys Thr Lys Glu Ser Leu Asp Val Ser

430 435 440 445

aga ctg acc cgg gaa ggt ggt ccc ctg ctg tat gaa ggc atc agt ctc 1394

Arg Leu Thr Arg Glu Gly Gly Pro Leu Leu Tyr Glu Gly Ile Ser Leu

450

455

460

acc atg aac tcc aaa gtc ttg aat ggc tcc cag cgg gtg gtg atg gat 1442

Thr Met Asn Ser Lys Val Leu Asn Gly Ser Gln Arg Val Val Met Asp
465
470
475

ggt gtg atc tct gat gat gag tgc cag gag ctg cag aga ctg acc aat 1490 Gly Val Ile Ser Asp Asp Glu Cys Gln Glu Leu Gln Arg Leu Thr Asn 480 485 490

gcg gca gca act tcg gga gat ggc tac cga ggt cag acc tcc cca cac 1538

Ala Ala Ala Thr Ser Gly Asp Gly Tyr Arg Gly Gln Thr Ser Pro His

495 500 505

acc cca aat gaa aag ttc tat ggt gtt act gtc ctc aaa gct ctc aag 1586

Thr Pro Asn Glu Lys Phe Tyr Gly Val Thr Val Leu Lys Ala Leu Lys

510 525

ctc ggg cag gaa gga aaa gtt cct ctg cag agt gcc cgc atg tac tac 1634
Leu Gly Gln Glu Gly Lys Val Pro Leu Gln Ser Ala Arg Met Tyr Tyr
530 535 540

aac gtg aca gag aag gtg cgg cgc gtc atg gag tcc tac ttc cgc ctg 1682
Asn Val Thr Glu Lys Val Arg Arg Val Met Glu Ser Tyr Phe Arg Leu
545 550 555

gac acg ccc ctc tat ttc tct tat tcc cac ttc gtg tgc cgc act gca 1730

Asp Thr Pro Leu Tyr Phe Ser Tyr Ser His Phe Val Cys Arg Thr Ala

560 565 570

ata gaa gag tca cag gct gag agg aag gac agt agc cac ccc gtc cac 1778

Ile Glu Glu Ser Gln Ala Glu Arg Lys Asp Ser Ser His Pro Val His

575 580 585

gtg gat aac tgc atc ctg aat gcc gaa gcc ttc atg tgt atc aag gag 1826 Val Asp Asn Cys Ile Leu Asn Ala Glu Ala Phe Met Cys Ile Lys Glu 590 595 600 605

ccc cca gca tac acg ttc cgg gaa tac agc gcc atc ctt tac ctc aat 1874
Pro Pro Ala Tyr Thr Phe Arg Glu Tyr Ser Ala Ile Leu Tyr Leu Asn
610 615 620

ggc gac ttc gat gga gga aac ttt tac ttc aca gaa cta gat gcc aag 1922

Gly Asp Phe Asp Gly Gly Asn Phe Tyr Phe Thr Glu Leu Asp Ala Lys
625 630 - 635

act gtg acg gca gag gtg cag ccc cag tgt gga agg gct gtg gga ttc 1970

Thr Val Thr Ala Glu Val Gln Pro Gln Cys Gly Arg Ala Val Gly Phe
640 645 650

tct tct ggc act gag aac cca cat gga gtg aag gct gtc acc agg ggg 2018

Ser Ser Gly Thr Glu Asn Pro His Gly Val Lys Ala Val Thr Arg Gly

655 660 665

cag cgc tgc gcc atc gcc ctg tgg ttc acg ctg gat cct cgg cac agt 2066

Gln Arg Cys Ala Ile Ala Leu Trp Phe Thr Leu Asp Pro Arg His Ser

670 680 685

gag aga gac agg gtg cag gca gat gac ctg gtg aag atg ctg ttc agc 2114

Glu Arg Asp Arg Val Gln Ala Asp Asp Leu Val Lys Met Leu Phe Ser

690 695 700

cca gaa gag gtg gac ctc ccc cag gaa cag ccc ctg cct gac cag cag 2162
Pro Glu Glu Val Asp Leu Pro Gln Glu Gln Pro Leu Pro Asp Gln Gln
705 710 715

ggt tcg cca gag cct gga gaa gag ttt ctg cat ggt gct act gtt ctt 2210 Gly Ser Pro Glu Pro Gly Glu Glu Phe Leu His Gly Ala Thr Val Leu 720 725 730 gga gtg ggc ata gca gga cac act ctt ctc tgg gct tgg ctg
Gly Val Gly Ile Ala Gly His Thr Leu Leu Trp Ala Trp Leu
735 740 745

taggeteaga atgeaggeee agaaceaeee tggggeetat gtaggeaget geegteagea 2312

gcgtgatata tttaagtgtc tgtaaagaca accaaagaat aaatgatttg tgtttttaaa 2372

aagnaaaaaa aaaaaaaaat taaaaatttg cgcggccgca agaa

2416

2252

<210> 6

<211> 747

<212> PRT

<213> Mus musculus

<400> 6

Met Ala Val Thr Lys Gly Gly Cys Trp His Asp Ala Ser Gly Arg Arg

1 5 10 15

Arg Arg Arg Leu Thr Gly Cys Gly Glu Ser Glu Pro Gly Trp Asp Val
20 25 30

Ala Ala Pro Asp Leu Leu Tyr Ala Glu Gly Thr Ala Ala Tyr Ser Arg

35 40 45

Arg Asp Trp Pro Gly Val Val Leu Asn Met Glu Arg Ala Leu Arg Ser 50 55 60

Arg Ala Ala Leu Arg Ala Leu Arg Leu Arg Cys Arg Thr Arg Cys Ala
65 70 75 80

Thr Glu Leu Pro Trp Ala Pro Asp Leu Asp Leu Gly Pro Asp Pro Ser

85 90 95

Leu Ser Gln Asp Pro Gly Ala Ala Ala Leu His Asp Leu Arg Phe Phe
100 105 110

Gly Ala Val Leu Arg Arg Ala Ala Cys Leu Arg Arg Cys Leu Gly Pro 115 120 125

Pro Ser Ala His Leu Leu Ser Glu Glu Leu Asp Leu Glu Phe Asn Lys
130 135 140

Arg Ser Pro Tyr Asn Tyr Leu Gln Val Ala Tyr Phe Lys Ile Asn Lys
145 150 155 160

Leu Glu Lys Ala Val Ala Ala Ala His Thr Phe Phe Val Gly Asn Pro 165 170 175

Glu His Met Glu Met Arg Gln Asn Leu Asp Tyr Tyr Gln Thr Met Ser

The state that the color and the state of th

Gly Val Lys Glu Ala Asp Phe Arg Asp Leu Glu Ala Lys Pro His Met

His Glu Phe Arg Leu Gly Val Arg Leu Tyr Ser Glu Glu Lys Pro Gln

Glu Ala Val Pro His Leu Glu Ala Ala Leu Gln Glu Tyr Phe Val Ala

Asp Glu Glu Cys Arg Ala Leu Cys Glu Gly Pro Tyr Asp Tyr Asp Gly

Tyr Asn Tyr Leu Asp Tyr Ser Ala Asp Leu Phe Gln Ala Ile Thr Asp

His Tyr Val Gln Val Leu Asn Cys Lys Gln Asn Cys Val Thr Glu Leu

Ala Ser His Pro Ser Arg Glu Lys Pro Phe Glu Asp Phe Leu Pro Ser

His Tyr Asn Tyr Leu Gln Phe Ala Tyr Tyr Asn Ile Gly Asn Tyr Thr Gln Ala Ile Glu Cys Ala Lys Thr Tyr Leu Leu Phe Phe Pro Asn Asp 325 330 335

Glu Val Met His Gln Asn Leu Ala Tyr Tyr Thr Ala Met Leu Gly Glu

340 345 350

Glu Glu Ala Ser Ser Ile Ser Pro Arg Glu Asn Ala Glu Glu Tyr Arg 355 360 365

Arg Pro Asn Leu Leu Glu Lys Glu Leu Leu Phe Phe Ala Tyr Asp Ile 370 375 380

Phe Gly Ile Pro Phe Val Asp Pro Asp Ser Trp Thr Pro Glu Glu Val 385 390 395 400

Ile Pro Lys Arg Leu Gln Glu Lys Gln Lys Ser Glu Arg Glu Thr Ala
405 410 415

Val Arg Ile Ser Gln Glu Ile Gly Asn Leu Met Lys Glu Ile Glu Thr
420 425 430

Leu Val Glu Glu Lys Thr Lys Glu Ser Leu Asp Val Ser Arg Leu Thr
435 440 445

Arg Glu Gly Gly Pro Leu Leu Tyr Glu Gly Ile Ser Leu Thr Met Asn 450 455 460 Ser Lys Val Leu Asn Gly Ser Gln Arg Val Val Met Asp Gly Val Ile 465 470 475 480

Ser Asp Asp Glu Cys Gln Glu Leu Gln Arg Leu Thr Asn Ala Ala Ala 485 490 495

Thr Ser Gly Asp Gly Tyr Arg Gly Gln Thr Ser Pro His Thr Pro Asn
500 505 510

Glu Lys Phe Tyr Gly Val Thr Val Leu Lys Ala Leu Lys Leu Gly Gln
515 520 525

Glu Gly Lys Val Pro Leu Gln Ser Ala Arg Met Tyr Tyr Asn Val Thr
530 535 540

Glu Lys Val Arg Arg Val Met Glu Ser Tyr Phe Arg Leu Asp Thr Pro 545 550 555 560

Leu Tyr Phe Ser Tyr Ser His Phe Val Cys Arg Thr Ala Ile Glu Glu
565 570 575

Ser Gln Ala Glu Arg Lys Asp Ser Ser His Pro Val His Val Asp Asn 580 585 590

Cys Ile Leu Asn Ala Glu Ala Phe Met Cys Ile Lys Glu Pro Pro Ala

Tyr Thr Phe Arg Glu Tyr Ser Ala Ile Leu Tyr Leu Asn Gly Asp Phe

Asp Gly Gly Asn Phe Tyr Phe Thr Glu Leu Asp Ala Lys Thr Val Thr

Ala Glu Val Gln Pro Gln Cys Gly Arg Ala Val Gly Phe Ser Ser Gly

Thr Glu Asn Pro His Gly Val Lys Ala Val Thr Arg Gly Gln Arg Cys

Ala Ile Ala Leu Trp Phe Thr Leu Asp Pro Arg His Ser Glu Arg Asp

Arg Val Gln Ala Asp Asp Leu Val Lys Met Leu Phe Ser Pro Glu Glu

Val Asp Leu Pro Gln Glu Gln Pro Leu Pro Asp Gln Gln Gly Ser Pro

Glu Pro Gly Glu Glu Phe Leu His Gly Ala Thr Val Leu Gly Val Gly

Ile Ala Gly His Thr Leu Leu Trp Ala Trp Leu

740 745

<210> 7

<211> 2322

<212> DNA

<213> Mus musculus

<220>

<221> CDS

<222> (12).. (1637)

1

<400> 7

ggagcaaggc c atg gcg gtg acg aaa gga ggc tgc tgg cac gat gct agc 50

Met Ala Val Thr Lys Gly Gly Cys Trp His Asp Ala Ser

10

5

ggt cgc cgc cgc cgc ctt acg ggt tgc ggc gag tct gag ccg gga 98

Gly Arg Arg Arg Arg Leu Thr Gly Cys Gly Glu Ser Glu Pro Gly

20 25

tgg gac gtg gca gcc cct gac ctg ctt tac gca gag ggg acc gcg gcc 146
Trp Asp Val Ala Ala Pro Asp Leu Leu Tyr Ala Glu Gly Thr Ala Ala
30 35 40 45

tac	tcg	cgc	agg	gac	tgg	ccc	ggg	gtg	gtc	ctg	aac	atg	gag	cgg	gct	194
Tyr	Ser	Arg	Arg	Asp	Trp	Pro	Gly	Val	Val	Leu	Asn	Met	Glu	Arg	Ala	
				50					55					60		
ctg	cgc	tcg	cgg	gcg	gcc	ctg	cgt	gcc	ctc	cgc	ctg	cgc	tgc	cgc	aca	242
Leu	Arg	Ser	Arg	Ala	Ala	Leu	Arg	Ala	Leu	Arg	Leu	Arg	Cys	Arg	Thr	
			65					70					75			
cgc	tgt	gcc	acc	gaa	ctg	ccg	tgg	gca	ccg	gac	ctg	gat	ctc	ggt	ccg	290
Arg	Cys	Ala	Thr	Glu	Leu	Pro	Trp	Ala	Pro	Asp	Leu	Asp	Leu	Gly	Pro	
		80					85					90				
gac	ccc	agc	ctg	agc	cag	gac	ccg	ggc	gcc	gcc	gcc	ctg	cac	gac	ctg	338
Asp	Pro	Ser	Leu	Ser	Gln	Asp	Pro	Gly	Ala	Ala	Ala	Leu	His	Asp	Leu	
	95					100					105					
cgc	ttc	ttc	gga	gcc	gtg	ctg	cgc	cgt	gcc	gcc	tgc	cta	cgc	cgc	tgc	386
Arg	Phe	Phe	Gly	Ala	Val	Leu	Arg	Arg	Ala	Ala	Cys	Leu	Arg	Arg	Cys	
110					115					120					125	
ctc	ggg	ccg	ccc	tct	gcc	cac	ttg	ctg	agt	gaa	gaa	ctg	gac	ctg	gag	434
Leu	Gly	Pro	Pro	Ser	Ala	His	Leu	Leu	Ser	Glu	Glu	Leu	Asp	Leu	Glu	
				130					135					140		

ttc aac aag cgg agc ccg tac aac tac ctg cag gtc gcc tat ttc aag 482 Phe Asn Lys Arg Ser Pro Tyr Asn Tyr Leu Gln Val Ala Tyr Phe Lys and a strong to the strong terms and strong terms and strong terms and the strong terms are strong terms are strong terms and the strong terms are strong terms are strong terms and the strong terms are strong terms and the strong terms are str

											-					
ata	aac	aag	ctg	gag	aaa	gct	gtg	gct	gcg	gca	cac	acc	ttc	ttt	gtg	530
Ile	Asn	Lys	Leu	Glu	Lys	Ala	Val	Ala	Ala	Ala	His	Thr	Phe	Phe	Val	
		160					165					170				
ggc	aat	cct	gag	cac	atg	gag	atg	cgg	cag	aac	ctc	gac	tat	tac	caa	578
Gly	Asn	Pro	Glu	His	Met	Glu	Met	Arg	Gln	Asn	Leu	Asp	Tyr	Tyr	Gln	
	175					180					185					

acc atg tct ggg gtg aag gag gca gac ttc agg gat ctc gag gcc aag 626
Thr Met Ser Gly Val Lys Glu Ala Asp Phe Arg Asp Leu Glu Ala Lys
190 195 200 205

ccc cat atg cat gag ttt cgg ctg ggg gta cga ctc tac tca gag gag 674

Pro His Met His Glu Phe Arg Leu Gly Val Arg Leu Tyr Ser Glu Glu
210 215 220

aag cca cag gaa gct gtg ccc cac ctg gag gcg gca ctg caa gag tac 722

Lys Pro Gln Glu Ala Val Pro His Leu Glu Ala Ala Leu Gln Glu Tyr
225 230 235

ttt gtg gcc gat gag gag tgc cgt gcc ctc tgc gaa ggg ccc tat gac 770

Phe Val Ala Asp Glu Glu Cys Arg Ala Leu Cys Glu Gly Pro Tyr Asp

240 245 250

tac	gac	ggc	tac	aac	tac	cta	gac	tac	agc	gct	gac	ctc	ttc	cag	gcc	818
Tyr	Asp	Gly	Tyr	Asn	Tyr	Leu	Asp	Tyr	Ser	Ala	Asp	Leu	Phe	Gln	Ala	
	255					260					265					
atc	aca	gat	cat	tac	gtc	cag	gtc	ctc	aac	tgt	aag	cag	aac	tgt	gtc	866
Ile	Thr	Asp	His	Tyr	Val	Gln	Val	Leu	Asn	Cys	Lys	Gln	Asn	Cys	Val	
270					275					280					285	
acg	gag	ctg	gct	tcc	cac	cca	agt	agg	gaa	aag	ccc	ttt	gaa	gac	ttc	914
Thr	Glu	Leu	Ala	Ser	His	Pro	Ser	Arg	Glu	Lys	Pro	Phe	Glu	Asp	Phe	
				290					295					300		
ctc	cct	tca	cac	tat	aat	tac	cta	cag	ttt	gcc	tac	tac	aac	att	ggg	962
Leu	Pro	Ser	His	Tyr	Asn	Tyr	Leu	Gln	Phe	Ala	Tyr	Tyr	Asn	Ile	Gly	
			305					310					315			
aac	tat	aca	caa	gct	att	gaa	tgt	gcc	aag	acc	tac	ctc	ctc	ttc	ttt	1010
Asn	Tyr	Thr	Gln	Ala	Ile	Glu	Cys	Ala	Lys	Thr	Tyr	Leu	Leu	Phe	Phe	
`		320					325					330				
ccc	aat	gat	gag	gtg	atg	cac	cag	aat	ctg	gct	tat	tac	aca	gcc	atg	1058
Pro	Asn	Asp	Glu	Val	Met	His	Gln	Asn	Leu	Ala	Tyr	Tyr	Thr	Ala	Met	
	335					340					345					

ctt gga gaa gaa gag gcc agc tcc atc agc ccc agg gag aat gcc gag 1106 Leu Gly Glu Glu Glu Ala Ser Ser Ile Ser Pro Arg Glu Asn Ala Glu

											-					
gaa	tac	cga	cgt	cca	aac	ctg	ttg	gag	aaa	gaa	ctg	ctt	ttc	ttc	gct	1154
Glu	Tyr	Arg	Arg	Pro	Asn	Leu	Leu	Glu	Lys	Glu	Leu	Leu	Phe	Phe	Ala	
				370					375					380		
tat	gac	att	ttt	gga	att	ccc	ttt	gtg	gat	ccc	gat	tca	tgg	act	cca	1202
Tyr	Asp	Ile	Phe	Gly	Ile	Pro	Phe	Val	Asp	Pro	Asp	Ser	Trp	Thr	Pro	
			385					390					395			
gaa	gaa	gtg	att	ccc	aag	aga	ttg	caa	gag	aaa	cag	aag	tct	gaa	cgg	1250
Glu	Glu	Val	Ile	Pro	Lys	Arg	Leu	Gln	Glu	Lys	Gln	Lys	Ser	Glu	Arg	
		400					405					410				
gaa	aca	gcc	gta	cgc	atc	tcc	cag	gag	att	ggg	aac	ctt	atg	aag	gaa	1298
Glu	Thr	Ala	Val	Arg	Ile	Ser	Gln	Glu	Ile	Gly	Asn	Leu	Met	Lys	Glu	
	415					420					425					
atc	gag	acc	ctt	gtg	gaa	gag	aag	acc	aag	gag	tct	ctg	gat	gtg	agc	1346
Ile	Glu	Thr	Leu	Val	Glu	Glu	Lys	Thr	Lys	Glu	Ser	Leu	Asp	Val	Ser	
430					435					440					445	
aga	ctg	acc	cgg	gaa	ggt	ggt	ссс	ctg	ctg	tat	gaa	ggc	atc	agt	ctc	1394
Arg	Leu	Thr	Arg	Glu	Gly	Gly	Pro	Leu	Leu	Tyr	Glu	Gly	Ile	Ser	Leu	
				450					455					460		

acc atg aac tcc aaa gtc ttg aat ggc tcc cag cgg gtg gtg atg gat 1442

Thr Met Asn Ser Lys Val Leu Asn Gly Ser Gln Arg Val Val Met Asp
465
470
475

ggt gtg atc tct gat gat gag tgc cag gag ctg cag aga ctg acc aat 1490 Gly Val Ile Ser Asp Asp Glu Cys Gln Glu Leu Gln Arg Leu Thr Asn 480 485 490

gcg gca gca act tcg gga gat ggc tac cga ggt cag acc tcc cca cac 1538

Ala Ala Ala Thr Ser Gly Asp Gly Tyr Arg Gly Gln Thr Ser Pro His

495 500 505

acc cca aat gaa aag ttc tat ggt gtt act gtc ctc aaa gct ctc aag 1586

Thr Pro Asn Glu Lys Phe Tyr Gly Val Thr Val Leu Lys Ala Leu Lys

510 525

ctc ggg cag gaa gga aaa gtt cct ctg cag agt gcc cgc acc gca ctg 1634
Leu Gly Gln Glu Gly Lys Val Pro Leu Gln Ser Ala Arg Thr Ala Leu
530 535 540

caa tagaagagte acaggetgag aggaaggaca gtagecacce egtecacgtg 1687 Gln

gataactgca teetgaatge egaageette atgtgtatea aggageeece agcatacaeg 1747

ttccgggaat acagcgccat cctttacctc aatggcgact tcgatggagg aaacttttac 1807

ttcacagaac tagatgccaa gactgtgacg gcagaggtgc agccccagtg tggaagggct 1867 gtgggattet ettetggeae tgagaaceea eatggagtga aggetgteae eagggggeag 1927 cgctgcgcca tcgccctgtg gttcacgctg gatcctcggc acagtgagag agacagggtg 1987 caggcagatg acctggtgaa gatgctgttc agcccagaag aggtggacct cccccaggaa 2047 cagcccctgc ctgaccagca gggttcgcca gagcctggag aagagtttct gcatggtgct 2107 actgttcttg gagtgggcat agcaggacac actcttctct gggcttggct gtaggctcag 2167 aatgcaggcc cagaaccacc ctggggccta tgtaggcagc tgccgtcagc agcgtgatat 2227 atttaagtgt ctgtaaagac aaccaaagaa taaatgattt gtgtttttaa aaagnaaaaa 2287 2322 aaaaaaaaaa ttaaaaattt gcgcggccgc aagaa

<210> 8

<211> 542

<212> PRT

<213> Mus musculus

Met Ala Val Thr Lys Gly Gly Cys Trp His Asp Ala Ser Gly Arg Arg

1 5 10 - 15

Arg Arg Arg Leu Thr Gly Cys Gly Glu Ser Glu Pro Gly Trp Asp Val
20 25 30

Ala Ala Pro Asp Leu Leu Tyr Ala Glu Gly Thr Ala Ala Tyr Ser Arg

35 40 45

Arg Asp Trp Pro Gly Val Val Leu Asn Met Glu Arg Ala Leu Arg Ser 50 55 60

Arg Ala Ala Leu Arg Ala Leu Arg Leu Arg Cys Arg Thr Arg Cys Ala
65 70 75 80

Thr Glu Leu Pro Trp Ala Pro Asp Leu Asp Leu Gly Pro Asp Pro Ser

85 90 95

Leu Ser Gln Asp Pro Gly Ala Ala Ala Leu His Asp Leu Arg Phe Phe
100 105 110

Gly Ala Val Leu Arg Arg Ala Ala Cys Leu Arg Arg Cys Leu Gly Pro 115 120 125

Pro Ser Ala His Leu Leu Ser Glu Glu Leu Asp Leu Glu Phe Asn Lys

130 135 140

Arg Ser Pro Tyr Asn Tyr Leu Gln Val Ala Tyr Phe Lys Ile Asn Lys

145 150 155 160

Leu Glu Lys Ala Val Ala Ala Ala His Thr Phe Phe Val Gly Asn Pro

165 170 175

Glu His Met Glu Met Arg Gln Asn Leu Asp Tyr Tyr Gln Thr Met Ser 180 185 190

Gly Val Lys Glu Ala Asp Phe Arg Asp Leu Glu Ala Lys Pro His Met

195 200 205

His Glu Phe Arg Leu Gly Val Arg Leu Tyr Ser Glu Glu Lys Pro Gln 210 215 220

Glu Ala Val Pro His Leu Glu Ala Ala Leu Gln Glu Tyr Phe Val Ala 225 230 235 240

Asp Glu Glu Cys Arg Ala Leu Cys Glu Gly Pro Tyr Asp Tyr Asp Gly

245 250 255

Tyr Asn Tyr Leu Asp Tyr Ser Ala Asp Leu Phe Gln Ala Ile Thr Asp
260 265 270

His Tyr Val Gln Val Leu Asn Cys Lys Gln Asn Cys Val Thr Glu Leu

											-				
Ala	Ser	His	Pro	Ser	Arg	Glu	Lys	Pro	Phe	G1u	Asp	Phe	Leu	Pro	Ser
	290					295					300				
His	Tyr	Asn	Tyr	Leu	Gln	Phe	Ala	Tyr	Tyr	Asn	Ile	Gly	Asn	Tyr	Thr
305					310					315					320
Gln	Ala	Ile	Glu	Cys	Ala	Lys	Thr	Tyr	Leu	Leu	Phe	Phe	Pro	Asn	Asp
				325					330					335	
Glu	Val	Met	His	Gln	Asn	Leu	Ala	Tyr	Tyr	Thr	Ala	Met	Leu	Gly	Glu
			340					345					350		
Glu	Glu	Ala	Ser	Ser	Ile	Ser	Pro	Arg	Glu	Asn	Ala	Glu	Glu	Tyr	Arg
		355					360				~	365			
Arg	Pro	Asn	Leu	Leu	Glu	Lys	Glu	Leu	Leu	Phe	Phe	Ala	Tyr	Asp	Ile
	370					375					380				

Phe Gly Ile Pro Phe Val Asp Pro Asp Ser Trp Thr Pro Glu Glu Val 385 390 395 400

Ile Pro Lys Arg Leu Gln Glu Lys Gln Lys Ser Glu Arg Glu Thr Ala
405 410 415

Val Arg Ile Ser Gln Glu Ile Gly Asn Leu Met Lys Glu Ile Glu Thr
420 425 430

Leu Val Glu Glu Lys Thr Lys Glu Ser Leu Asp Val Ser Arg Leu Thr
435 440 445

Arg Glu Gly Gly Pro Leu Leu Tyr Glu Gly Ile Ser Leu Thr Met Asn 450 455 460

Ser Lys Val Leu Asn Gly Ser Gln Arg Val Val Met Asp Gly Val Ile 465 470 475 480

Ser Asp Asp Glu Cys Gln Glu Leu Gln Arg Leu Thr Asn Ala Ala Ala 485 490 495

Thr Ser Gly Asp Gly Tyr Arg Gly Gln Thr Ser Pro His Thr Pro Asn
500 505 510

Glu Lys Phe Tyr Gly Val Thr Val Leu Lys Ala Leu Lys Leu Gly Gln
515 520 525

Glu Gly Lys Val Pro Leu Gln Ser Ala Arg Thr Ala Leu Gln
530 535 540

```
⟨211⟩ 28
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Artificially
      Synthesized Primer Sequence
<400> 9
ggatccaagg agcgggctct gcgctcgc
⟨210⟩ 10
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Artificially
```

Synthesized Primer Sequence

<400> 10

ccaagcttgg ctgtgtaata a

21

<211> 19 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Artificially Synthesized Primer Sequence <400> 11 tcattacatc caggtcctc <210> 12 <211> 20 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Artificially Symthesized Primer Sequence <400> 12

19

<400> 12

tttggagttc atggtgagac 20

```
<211> 38
<212> DNA
<213> Art
<220>
<223> Desc
```

<213> Artificial Sequence

<400> 13

agatctagat ctatggcggt acgcgcgttg aagctgct

38

<210> 14

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<400> 14

gtcgacgtcg acttcatagc tcatccttgg gcttcgatt

⟨211⟩ 40

<212> DNA

<213> Artificial Sequence

<220>

<400> 15

gtcgacgtcg actctaggtg ccctgctcac gggggccgat